Multimedia Appendix 2. Pseudo code of symptom frequencies based on vector distance (SF-DIST) and cosine similarity (SF-COS)

```
SF-COS :
Input : patient_symptoms (10x1), symptoms_frequencies (10x4)
if all patient_symptoms == 0:
    Return LOW RISK

For each column of symptoms_frequencies:
        similarity = cosine_similarity(patient_symptoms, symptoms_frequencies[column])
similarity = normalise(similarity)
similarity[COVID_19] = similarity[COVID_19]*(Area_Risk_Factor + Contact_Risk_Factor)
Output : return disease with maximum similarity
```

```
SF-DIST:
Input : patient_symptoms (10x1), symptoms_frequencies (10x4)

if all patient_symptoms == 0:
    Return LOW RISK

For each column of symptoms_frequencies:
    similarity = sum(abs((patient_symptoms - symptoms_frequencies[column]))

similarity = normalise(similarity)

similarity[COVID_19] = similarity[COVID_19]*(Area_Risk_Factor + Contact_Risk_Factor)

Output : return disease with maximum similarity
```